1653

RAW SEQUENCE LISTING

PATENT APPLICATION: US/08/962,560B

DATE: 09/14/2000

TIME: 17:33:46

Input Set : A:\PTO.txt

Output Set: N:\CRF3\09142000\H962560B.raw

Does Not Comply Corrected Diskette Needed 5ec PP.

```
4 <110> APPLICANT: Hilton, Douglas J.
        Alexander, Warren S.
        Viney, Elizabeth M.
        Wilson, Tracy A.
        Richardson, Rachel
        Starr, Robyn
        Nicholson, Sandra E.
        Metcalf, Donald
        Nicola, Nicos A
14 <120> TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC AGENTS
16 <130> FILE REFERENCE: Davies Collison Cave
18 <140> CURRENT APPLICATION NUMBER: 08/962,560B
19 <141> CURRENT FILING DATE: 1997-10-31
21 <160> NUMBER OF SEQ ID NOS: 68
23 <170> SOFTWARE: PatentIn Ver. 2.1
```

ERRORED SEQUENCES

```
137 <210> SEO ID NO: 4
     138 <211> LENGTH: 212
     139 <212> TYPE: PRT
     140 <213> ORGANISM: Mus musculus
     142 <400> SEQUENCE: 4
     143 Met Val Ala Arg Asn Gln Val Ala Ala Asp Asn Ala Ile Ser Pro Ala 144 \phantom{0} 1 5 \phantom{0} 10 \phantom{0} 15
     146 Ala Glu Pro Arg Arg Ser Glu Pro Ser Ser Ser Ser Ser Ser Ser
     149 Ser Pro Ala Ala Pro Val Arg Pro Arg Pro Cys Pro Ala Val Pro Ala
150 35 40 45
     152 Pro Ala Pro Gly Asp Thr His Phe Arg Thr Phe Arg Ser His Ser Asp 50 55 60
     155 Tyr Arg Arg Ile Thr Arg Thr Ser Ala Leu Leu Asp Ala Cys Gly Phe 156 \, 65 \, 70 \, 75 \, 80
     158 Tyr Trp Gly Pro Leu Ser Val His Gly Ala His Glu Arg Leu Arg Ala
159
85
90
95
E--> 161 Glu Pro Va Cly Thr Phe Leu Val Arg Asp Ser Arg Gln Arg Asn Cys
162 100 110
110 110 Arg Asp Ser Cly Pro Thr Ser Ile Arg
     164 Phe Phe Ala Leu Ser Val Lys Met Ala Ser Gly Pro Thr Ser Ile Arg
165 115 120 125
     167 Val His Phe Gln Ala Gly Arg Phe His Leu Asp Gly Ser Arg Glu Thr
168 130 135 140
     171 Phe Asp Cys Leu Phe Glu Leu Leu Glu His Tyr Val Ala Ala Pro Arg
172 145 150 . 155 160
     177 Glu Leu Cys Arg Gln Arg Ile Val Ala Ala Val Gly Arg Glu Asn Leu
```

Invalid amino acid designator.

```
Input Set : A:\PTO.txt
                      Output Set: N:\CRF3\09142000\H962560B.raw
                                           185
     180 Ala Arg Ile Pro Leu Asn Pro Val Leu Arg Asp Tyr Leu Ser Ser Phe
                                       200
                 195
     183 Pro Phe Gln Ile
             210
     1644 <210> SEQ ID NO: 24
     1645 <211> LENGTH: 2019
     1646 <212> TYPE: DNA
     1647 <213> ORGANISM: Mus musculus
     1649 <220> FEATURE:
     1650 <221> NAME/KEY: UNSURE
     1651 <222> LOCATION: (1981)
     1652 <223> OTHER INFORMATION: N is unsure
     1654 <220> FEATURE:
     1655 <221> NAME/KEY: UNSURE
     1656 <222> LOCATION: (1992)
     1657 <223> OTHER INFORMATION: N is unsure
     1659 <220> FEATURE:
                                         -> n is at location 2000
     1660 <221> NAME/KEY: UNSURE
1661 <222> LOCATION (2010)
     1662 <223> OTHER INFORMATION: N is unsure
     1664 <400> SEQUENCE: 24
     1665 ggcacgaggc ggggtcaggg cggaggctga ggaccaagta ggcatggcgg agggcgggac 60
     1667 cggccccgat ggacgggccg gcccgggacc cgcaggtcct aatctgaagg agtggctgag 120
1669 ggagcagttc tgtgaccatc cactggagca ctgtgacgat acaagactcc atgatgcagc 180
     1671 ctatgtaggg gacctccaga ccctcaggaa cctactgcaa gaggaggact accggagccg 240
1673 catcaatgag aagtetgtet ggtgetgegg etggetteee tgcacaceae tgaggatege 300
     1675 agccactgca ggccatggga actgtgtgga cttcctcata cgcaaagggg ccgaggtgga 360
     1677 cctggtggat gtcaaggggc agactgcct gtatgtggct gtagtgaacg ggcacttgga 420
     1679 gagcactgag atcettttgg aagetggtge tgateecaae ggeageegge accaeegeag 480
     1681 cactectyty taccatycet ytegtytygy tagggacyae atectyaagg etettateag 540
     1683 gtatggggca gatgttgatg tcaaccatca totgaattot gacaccoggc cocetttttc 600
     1685 acggcggcta acctccttgg tggtctgtcc tctatacatc agtgctgcct accataacct 660
     1687 teagtgette aggetgetet tgeaggetgg ggeaaateet gactteaatt geaatggeee 720
     1689 tgtcaacacc caggagttct acaggggatc ccctgggtgt gtcatggatg ctgtcctgcg 780
     1691 ccatggctgt gaagcagcct tcgtgagtct gttggtagag tttggagcca acctgaacct 840
     1693 ggtgaagtgg gaatcoctgg gcccagaggc aagaggcaga agaaagatgg atcctgaggc 900 1695 cttgcaggtc tttaaagagg ccagaagtat tcccaggacc ttgctgagtt tgtgccgggt 960
     1697 ggctgtgaga agagetettg geaaataceg actgeatetg gttecetege tgeegetgee 1020
                                                                                             => should be 1500
     1699 agaccccata aagaagtttt tgctttatga gtagcattca catgcagtgc tgactgcaat 1080
     1701 gtggaageeg ateacetgea gtgaaaaetg acacagaete tggeateetg ggaaceatgg 1140
     1703 cctgtgctgc cagcttgatc cttggctgtc agtgaagaaa aaacggctgt gttctcttgg 1200
     1705 actgtgattc tatctcaggt gcttgggcca tcgaacgctc cttgagtcat tgtcaactga 1260
     1707 gaggcacata caaacttaat tttgttcctc ttcagtctct ctgttttgga ttcttcctgg 1320
     1709 caatgtgtgc agcatgggct gagcctggtg attgccctag tggggaaggc ttttttctcc 1380
     1711 aggctatgca totatttatg ttcctacttt gcaatttatt gttcttttaa ggcttgatat 1440
E--> 1713 caaaacagaa agaggtttgt taagaaaaga tatagggaga aaggaattcc ggttccgtgc (1550)
     1715 acttgctage etgettteet tgeetgggtt tgtetgteta tgetgeetgg tgeacatece 1560
     1717 ttetetttge tgccactgtt ctattttggg agttgtette egtetaagat ggettetggg 1620
```

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1719 gttctatctt attgcacaga ggtcccagaa cagtgttcat agggcaccat ctgctctgcc 1680 1721 aagggtttte tgatgtetta eettgaggat etteagacag tggttaeett taggagaeee 1740
1723 aeetggaaet aaceattaag tgaetgeeea eatteagate agggaeeate ttaatagtae 1800 1725 teactgocag teeteacaag agaagatgae aegggtgete tetteagaea eteccataea 1860 1727 ggaagttgga aaatgtettg gteacetggg ttgtteecag getacaactt ettggtgtte 1920 1729 cactaaracc agratatect agtittitigg gitigactigt cectececae titeetigaa 1980 W--> 1731 neceaatgee entitigikin gitigettee etaaaaktt 2019

1 is at location 2000, not 2010

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY DATE: 09/14/2000 PATENT APPLICATION: US/08/962,560B TIME: 17:33:47

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```
L:161 M:333 E: Wrong sequence grouping, Amino acids not in groups!
L:161 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:1
L:1085 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1117 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1137 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:1237 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1279 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18
L:1713 M:254 E: No. of Bases conflict, LENGTH:Input:1550 Counted:1500 SEQ:24
L:1731 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:1775 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:1916 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:1920 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:2524 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:41
L:2679 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3419 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3423 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3431 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3433 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3434 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
L:3437 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51
```